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(FILE 'HOME' ENTERED AT 15:27:02 ON 11 MAR 2008)

FILE 'USPATFULL, WPIDS' ENTERED AT 15:27:26 ON 11 MAR 2008

L1 109 FILE USPATFULL

L2 5 FILE WPIDS

TOTAL FOR ALL FILES

L3 114 S ((QUANTUM MECHANICS) AND (MOLECULAR MECHANICS) AND SIMULAT?

L4 85 FILE USPATFULL

L5 2 FILE WPIDS

TOTAL FOR ALL FILES

L6 87 S ((QUANTUM MECHANICS) AND (MOLECULAR MECHANICS) AND SIMULAT? AN

FILE 'USPATFULL' ENTERED AT 15:44:01 ON 11 MAR 2008

L7 63 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?

L8 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?

L9 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?

L10 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?

L11 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?

L12 86 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?

L13 48 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) (P) SIMULAT?

=> d 113 5,12,13 bib,kwic

L13 ANSWER 5 OF 48 USPATFULL on STN

AN 2007:49718 USPATFULL

TI Molecular simulation method and device

IN Yonezawa, Yasushige, Suita-shi, JAPAN

Takada, Toshikazu, Minato-ku, JAPAN

Nakata, Kazuto, Minato-ku, JAPAN

Sakuma, Toshihiro, Minato-ku, JAPAN

Nakamura, Haruki, Suita-shi, JAPAN

PA NEC CORPORATION, Tokyo, JAPAN (non-U.S. corporation)

OSAKA UNIVERSITY, Osaka, JAPAN (non-U.S. corporation)

PI US 2007043545 A1 20070222

AI US 2004-573023 A1 20040922 (10)

WO 2004-JP13808 20040922

20060322 PCT 371 date

PRAI JP 2003-329751 20030922

DT Utility

FS APPLICATION

LREP DICKSTEIN SHAPIRO LLP, 1177 AVENUE OF THE AMERICAS (6TH AVENUE), NEW

YORK, NY, 10036-2714, US

CLMN Number of Claims: 18

ECL Exemplary Claim: 1

DRWN 3 Drawing Page(s)

LN.CNT 754

SUMM The present invention relates to a method and a device for performing molecular simulation by a quantum chemical technique, and more particularly relates to a molecular simulation method and a device by a QM/MM (Quantum Mechanics/Molecular Mechanics) method in which the ab initio molecular orbital method and the molecular mechanics method are combined as one theoretical system among theoretical techniques of the quantum chemistry.

SUMM . . . the like are limited. Therefore, it is considered that only the sites and the vicinity thereof are subjected to accurate simulation about electron states. Accordingly, in a chemical system including many molecules, the QM/MM method is proposed in which a

molecule or a part of molecule is divided into a QM (Quantum Mechanics) space where a noted chemical phenomenon occurs and a secondary MM (Molecular Mechanics) space other than the QM space and in which the QM space is processed by a quantum mechanical scheme such as the ab initio molecular orbital method and the MM space is described as empirical potential such as molecular mechanics [1]. Advantages of this method are following:

L13 ANSWER 12 OF 48 USPATFULL on STN

AN 2005:319379 USPATFULL

TI Methods for molecular property modeling using virtual data

IN Duffy, Nigel P., San Francisco, CA, UNITED STATES

Lanza, Guido, San Francisco, CA, UNITED STATES

Yu, Jessen, San Francisco, CA, UNITED STATES

Mydlowec, William, San Francisco, CA, UNITED STATES

PI US 2005278124 A1 20051215

AI US 2005-74587 A1 20050308 (11)

PRAI US 2004-579619P 20040614 (60)

DT Utility

FS APPLICATION

LREP RAYMOND R. MOSER JR., ESQ., MOSER IP LAW GROUP, 1040 BROAD STREET, 2ND FLOOR, SHREWSBURY, NJ, 07702, US

CLMN Number of Claims: 39

ECL Exemplary Claim: 1

DRWN 4 Drawing Page(s)

LN.CNT 808

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

DETD . . . included in the training data may be provided using "virtual data," and may include information obtained from reasonable assumptions, computer simulations, or other modeling efforts. For example, computer simulations may be performed that simulate the physics of the molecular property of interest using molecular mechanics or quantum mechanics. Property information may also be obtained from laboratory experimentation or published literature sources. Additionally, property information may include a measure. . .

L13 ANSWER 13 OF 48 USPATFULL on STN

AN 2005:196237 USPATFULL

TI Lead molecule cross-reaction prediction and optimization system

IN Kita, David, Milpitas, CA, UNITED STATES

Fodor, Eniko, Fremont, CA, UNITED STATES

Prakash, Adityo, Fremont, CA, UNITED STATES

PA Verseon, Milpitas, CA, UNITED STATES (U.S. corporation)

PI US 2005170379 A1 20050804

AI US 2004-966341 A1 20041014 (10)

PRAI US 2003-511474P 20031014 (60)

DT Utility

FS APPLICATION

LREP TOWNSEND AND TOWNSEND AND CREW, LLP, TWO EMBARCADERO CENTER, EIGHTH FLOOR, SAN FRANCISCO, CA, 94111-3834, US

CLMN Number of Claims: 70

ECL Exemplary Claim: 1

DRWN 25 Drawing Page(s)

LN.CNT 3120

DETD . . . earlier in both the background section and the detailed technical description, including standard shape complementarity methods, conventional techniques based on molecular mechanics paradigms, molecular dynamics and/or quantum mechanics simulations, QSAR, free energy perturbation theory, or even the utilization of various empirically derived or knowledge-based scoring

function.

=> d his

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L9 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?  
L10 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?  
L11 0 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?  
L12 86 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) AND SIMULAT?  
L13 48 S (((QUANTUM MECHANICS) (P) (MOLECULAR MECHANICS)) (P) SIMULAT?

=> d l2 1-5 bib,kwic

YOU HAVE REQUESTED DATA FROM FILE 'WPIDS' - CONTINUE? (Y)/N:y

L2 ANSWER 1 OF 5 WPIDS COPYRIGHT 2008 THE THOMSON CORP on STN  
AN 2008-A36197 [03] WPIDS  
DNN N2008-028101 [03]  
TI Method of positioning polyphenols compound antioxidant point  
DC S03; T01  
IN LV L  
PA (UYNA-N) UNIV NANJING  
CYC 1  
PIA CN 101000354 A 20070718 (200803)\* ZH [1]  
ADT CN 101000354 A CN 2006-10166311 20061226  
PRAI CN 2006-10166311 20061226  
NOV NOVELTY - The invention claims a method of positioning polyphenols compound antioxidant point. specifically the MM + molecular mechanics method of testing polyphenols compound is used for processing primary optimizing structure calculation, then the semi-empirical AM1 quantum mechanics method is used for processing further optimizing calculation, finally Gaussian 03 program with abinitio calculation HF method is used for.. . as the amount of the net electronic charge of dissociation energy and atom, then the hyperoxidation revulsant is used for simulating oxidation reaction, the segregative oxidation metabolite is identified its structure, after that it assures the active point after oxidation reaction.. . only makes up the deficiency under the ideal condition calculating, but also avoids the inferential deviation of complex metabolite by simulating oxidation reaction, and it is able to test the polyphenols compound antioxidant point.

L2 ANSWER 2 OF 5 WPIDS COPYRIGHT 2008 THE THOMSON CORP on STN  
AN 2005-285012 [29] WPIDS

DNC C2005-088424 [29]  
 DNN N2005-233808 [29]  
 TI Molecular simulation in chemical industry, involves dividing  
 molecule into quantum mechanics space and  
 molecular mechanics space and applying non-empirical  
 molecule orbital method to quantum mechanics space  
 DC J04; T01  
 IN NAKAMURA H; NAKATA K; SAKUMA T; TAKADA T; YONEZAWA Y  
 PA (NIDE-C) NEC CORP; (OSAU-C) UNIV OSAKA  
 CYC 106  
 PIA WO 2005029385 A1 20050331 (200529)\* JA 28[5]  
 JP 2005514101 X 20061130 (200681) JA 18  
 US 20070043545 A1 20070222 (200717) EN  
 ADT WO 2005029385 A1 WO 2004-JP13808 20040922; JP 2005514101 X WO 2004-JP13808  
 20040922; JP 2005514101 X JP 2005-514101 20040922; US 20070043545 A1 WO  
 2004-JP13808 20040922; US 20070043545 A1 US 2006-573023 20060322  
 FDT JP 2005514101 X Based on WO 2005029385 A  
 PRAI JP 2003-329751 20030922  
 TI Molecular simulation in chemical industry, involves dividing  
 molecule into quantum mechanics space and  
 molecular mechanics space and applying non-empirical  
 molecule orbital method to quantum mechanics space  
 TT TT: MOLECULAR SIMULATE CHEMICAL INDUSTRIAL DIVIDE QUANTUM  
 MECHANICAL SPACE APPLY NON EMPIRICAL ORBIT METHOD  
 NOV NOVELTY - A molecule to be simulated is divided into  
 quantum mechanics (QM) space and molecular  
 mechanics (MM) space. A non-empirical molecule orbital method is  
 applied to the QM space with an empirical potential.  
 DETD DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the  
 following:  
 (1) a molecular simulation apparatus;  
 (2) a molecular simulation program; and  
 (3) recording media storing molecular simulation program.  
 USE  
 USE - Used for performing molecular simulation of a living  
 organism in the chemical industry, and during the manufacture of  
 pharmaceutical and functional food, and also in. . .  
 ADV ADVANTAGE - The molecular simulation is performed with high  
 reliability.  
 DRWD DESCRIPTION OF DRAWINGS - The figure shows a flowchart illustrating  
 molecular simulation. (Drawing includes non-English language  
 text).  
 L2 ANSWER 3 OF 5 WPIDS COPYRIGHT 2008 THE THOMSON CORP on STN  
 AN 2004-088951 [09] WPIDS  
 DNC C2004-036271 [09]  
 DNN N2004-071198 [09]  
 TI Development of molecular mechanics force field  
 parameters involves exporting the optimized force field parameters to  
 external molecular mechanics simulation  
 packages and saving optimize force field parameters to the database  
 DC B04; J04; T01  
 IN SUN H  
 PA (SUNH-I) SUN H  
 CYC 1  
 PIA US 20030195734 A1 20031016 (200409)\* EN 17[3]  
 US 6785665 B2 20040831 (200457) EN  
 ADT US 20030195734 A1 US 2002-139806 20020416  
 PRAI US 2002-139806 20020416  
 TI Development of molecular mechanics force field  
 parameters involves exporting the optimized force field parameters to

external molecular mechanics simulation  
 packages and saving optimize force field parameters to the database

TT TT: DEVELOP MOLECULAR MECHANICAL FORCE FIELD PARAMETER OPTIMUM EXTERNAL  
 SIMULATE PACKAGE SAVE DATABASE

NOV NOVELTY - Molecular mechanics force field parameters  
 are developed by importing molecular models representing the molecular  
 systems to be parameterized, validating the estimated and optimized force  
 field parameters, and exporting the optimized force field parameters in  
 required formats to external molecular mechanics  
 simulation packages and saving the molecular models, input data  
 and optimize force field parameters to the database.

DETD DETAILED DESCRIPTION - Development of molecular  
 mechanics force field parameters comprises creating or importing  
 molecular models that represent the molecular systems to be parameterized,  
 searching a database. . . the molecular models and stored molecular  
 models and retrieving stored parameters if complete matches are found,  
 preparing input data for quantum mechanics ab initio  
 calculations for the molecular models, importing calculated data of the  
 quantum mechanics ab initio calculations for the  
 molecular models, selecting force field type and functional forms, and  
 assigning atom types to the. . . set of mathematical formulas for the  
 molecular models, optimizing the initial force field parameters to fit the  
 input data of quantum mechanics ab initio  
 calculations, validating the optimized force field parameters, and  
 exporting the optimized force field parameters in required formats to  
 external molecular mechanics simulation  
 packages and saving the molecular models, input data and optimize force  
 field parameters to the database.

USE USE - For developing molecular mechanics force field  
 parameters for computer simulations of molecular systems, which  
 include molecules, clusters of molecules and clusters of atoms.

ADV ADVANTAGE - The inventive method is capable of rapidly and easily provides  
 high quality molecular mechanics force fields that are  
 required for successful computer modeling in chemical, pharmaceutical and  
 material industries

L2 ANSWER 4 OF 5 WPIDS COPYRIGHT 2008 THE THOMSON CORP on STN

AN 1999-418399 [35] WPIDS

DNC C1999-122896 [35]

DNN N1999-312326 [35]

TI Calculating relative stabilities of two molecules from conformational free  
 energies

DC A35; B02; B04; D15; D16; E37; T01

IN KOLOSSVARY I

PA (UYCO-C) UNIV COLUMBIA NEW YORK

CYC 23

PIA WO 9917222 A1 19990408 (199935)\* EN 88[4]  
 AU 9897796 A 19990423 (199935) EN  
 US 6178384 B1 20010123 (200107) EN

ADT WO 9917222 A1 WO 1998-US20368 19980929; US 6178384 B1 US 1997-940145  
 19970929; AU 9897796 A AU 1998-97796 19980920

FDT AU 9897796 A Based on WO 9917222 A

PRAI US 1997-940145 19970929

ADV. . . method should identify agents with greater specificity for, and  
 activity at, target sites. It does not require expensive free energy  
 simulations or computational 'alchemy'.

TECH. . . .  
 The low-energy minimum conformations are 15-35 kJ/mole above the lowest  
 energy conformation and the potential energy function is based on  
 molecular mechanics or on ab initio, semi-empirical or

density-functional quantum mechanics. The atomic co-ordinates used are external (Cartesian) or internal (bond lengths, bond or torsional angles) and Hi is exact or.. . .

Member(0003)

ABEQ US 6178384 . . .

method should identify agents with greater specificity for, and activity at, target sites. It does not require expensive free energy simulations or computational 'alchemy'.

L2 ANSWER 5 OF 5 WPIDS COPYRIGHT 2008 THE THOMSON CORP on STN

AN 1996-363502 [37] WPIDS

DNC C1996-114569 [37]

DNN N1996-306432 [37]

TI Construction of molecular models, e.g. for drug design - providing accurate 3-dimensional geometry by successive addition of fragments which regard the binding site region in a special way

DC B04; J04; S05; T01

IN FERINCZ J; KRIEG B

PA (FERI-I) FERINCZ J; (KRIE-I) KRIEG B

CYC 1

PIA DE 19504724 A1 19960808 (199637)\* DE 10[6]

ADT DE 19504724 A1 DE 1995-19504724 19950201

PRAI DE 1995-19504724 19950201

USE USE - The system is based on a numerical process (molecular mechanics, quantum mechanics, or molecular dynamics) or an expert system. It is useful for determining the 3-dimensional geometry of molecules in organic chemistry, biochemistry, molecular biology and related sciences, e.g. in understanding reaction mechanisms, developing and improving drugs ('drug design') and simulating chemical reactions using computers.